

**FIGURE 1a**  
(SEQ ID NO: 1 and SEQ ID NO: 2)

TTTAATCATG	GAATATTTCA	AACATACAGA	AAAATCACAG	AAAATAAATA	ACAACCACTC	ATTTATCTTC	-1101
TCCCCAACCC	CATGTAATAA	ATATTAAAT	ATTGTGTTAA	ATGCTAAATT	TAACACATGC	TAAAGGTTCC	-1031
TGGCTGGATG	TGGTGGCTCA	CGCCTGTAAT	CCCAGTACTT	TGGGAGGAGG	AGGTGGGAGG	ATTGCTTGAG	-961
TCCAGGAGCT	CGAGACCAGC	ATGGGCAACA	TAGTGCATC	TCGTCTCTAC	AAAAACAAA	AAAATTAGCT	-891
GGGCATGGTG	GTGTGCATCA	GTAATCCCAG	TGACTGGGAG	GCTGAGGTGG	GAGAATTGCT	TGAGTCTGGG	-821
AATTTGAGGC	TGCAGTGAGC	CCTGATCATG	CCACTGCATT	CCAGCATGGG	CGACATAGCA	AAACTTGTC	-751
AAAAAAAAAA	AAGTTTCCTC	TCTGCCCCAC	CATAGACAAC	CACTCTTCTG	ATTTCTATCT	TCGTAGATGA	-681
ATTTTGCCCA	TTCTCTTGTA	TATGAAAGGA	ACCAGACATT	AGGCATTCTG	GTGTCTGGTT	TCTTTCACCT	-611
AAGATAAAAT	TGAGTTAACC	TGTATTGTTG	TACAGAACTG	CAGTTTGTTT	TTTGTTATTT	ATTGTAAAGA	-541
CAGGGTCTGG	CTATGTTGCC	TAGGCTGGTC	TCGAACTGTT	GGCCTCAAGC	AATCCACCTG	CCAAGCTCTG	-471
GGACCACAGG	CATGAGCCAT	GGCATCTGAT	CKGTAGTTTG	ATCTTATTTT	TTGCTGAGTA	GTAGCCCATG	-401
GCATGACTTT	ATTATTTTGG	GTGTCCATTC	TCCTCTGGAG	GGGCTCTGCT	TTTTGAAACC	ACACCCTGGC	-331
CTAGCTCCCC	TTCTCCCTGC	CTCTCTGCAG	GCTCACATCC	ACATGCCAAG	ACCTCTGCAG	CCATTCTGCT	-261
TCCTGTCTTT	CCACTCCTGT	GGGACCTCAG	AGAGCTACGG	GGCTCCCTGG	GTACCAACTG	GCTCCTGAGG	-191
CCTGGGGGAG	GGTGGTCTTC	TGGGAGAAGG	AAGCCAGGTC	CCTGCAGGTT	GTGGAGGGGG	ACAGAATGAG	-121
GGTTTTTCCC	CAGGATGTTG	TTGGCCCCTG	CCCCCACTTC	TGTTCCATAA	TTAACCACGC	CCCTCCTACC	-51
CACTGTGCCC	CTCTTCCTGC	TGTGTGGAGG	CCCTGAATCA	TTATTTTAAC	TACCCCTGGG	GAGGGTGAGC	20
ACCTTCTGTG	CTCTGTCCCC	AACCTTCCAC	TTCCCCTCAA	CGCGCTGCTC	AGGCATGACC	TTCGGCACTG	90
					M T	P G T	

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FIGURE 1a (continued)

TGCTTCTTCT	GAGTG <u>g</u> taag	tggggccagg	gtgctgggga	gaagcttgga	ggagttctga	160 ggggactcca
V L L L S	caggctgggg	gctggtggtc	ggctccaacc	actcttatga	ggagctgagg	230 caggggagtg
cttcatgtgc	gagtggcccc	gagtcagtag	agtgtgacct	gaatgaagag	gggctcaggg	300 gctgtgctca
ggtggcgact	aagctacctc	tccagctggc	tatgttgtcc	caggcttccc	tgctcccact	370 catggagtcc
ctggtgtggg	tgacagaggt	ctccccagcc	tcccccgga	gtggaagcc	acagaagcca	440 ccagggaggg
ggaaagggtg	gacatcacct	ccctgggcct	nnnnnttccc	ccaagtctg	actgcacgta	510 gggaagaggc
ccctgctga	aaactgcatc	agagtcacat	INTRON 1 tcacgtgcca	tcaaaaatca	ggcttggtg	580 ggtgcggtg
ctcatgctta	taatcccagc	actttgggag	gccgagatgg	gcgtatcccc	tgaggtcagg	650 agtttgtgac
cagcctggcc	aacatgggtga	aaccccatct	ttaccaaaaa	tataaaaatt	agccgggcat	720 ggtggcgtgc
acttgtaatc	ccagctactt	gggaagctga	ggcaagagaa	tcgcttgaa	ccaggagacg	790 gaagttgcag
tgagctgaga	tcgtgccgtt	gcactccagc	ctcagcaaca	gagcgagact	ccatctcaaa	860 aaaaaaaaaa
aaaaaaagaa	aaaaaagaaa	aagaggctgg	gaggtcctag	ggattggggc	ttctttaact	930 cccagcctcc
ccgcccacca	aatattcctc	<u>ag</u> TCCTGGCT	TCTTATCATG	GATTCAACCT	GGATGTGGAG	1000 GAGCCTACGA
		V L A	S Y H	G F N L	D V E	E P T

**FIGURE 1b**  
**(SEQ ID NO: 1 and SEQ ID NO: 2)**

1070	TCTTCCAGGA I F Q E	GGATGCAGGC	GGCTTTGGGC	AGAGCGTGGT	GCAGTTCGGT	GGATCTCGgt	aggccccact
	INTRON 2 (3019 bp)						
4080	cccccaagtg	cccgctgctc	ccacccctcc	tgtggctgca	gtgacatggc	catggttgtg	tctccagACT <sub>L</sub>
4150	CGTGTGGGA V V G	GCACCCCTGG	AGGTGGTGGC	GGCCAACCCAG	ACGGGACGGC	TGTATGACTG	CGCAGCTGCC
4220	ACCGGCATGT	GCCAGCCCAT	CCCGCTGCAC	Agtgatgac	cacctgggaa	ttgggccccct	caacctcct
4290	T G M	C Q P I	P L H	INTRON 3			
4360	ggacccaact	gtgccccgcg	ttagcttcca	gtccagacct	tccccgcaaa	tgaagtgtgtg	ctgtgagtga
4430	gacccccggt	gtctgccctt	gcagtccgcg	CTGAGGCCGT	GAACATGTCC	TTGGGCCCTGA	CCCTGGCAGC
4500	CTCCACCAAC	GGCTCCCGGC	TCCTGgtgag	P E A V	N M S	L G L	T L A A
4570	S T N	G S R	L L	INTRON 4		ggggtgggggt	ggggcggggg
4640	gtgttgttgg	ggaggaggct	ggggctggga	gtgaaggagg	aggggctgct	agggactcct	ggctcacagg
	cttctgcctc	cagGCCTGTG	GCCCGACCCCT	GCACAGAGTC	TGTGGGGAGA	ACTCATACTC	AAAGGGTTCC
	TGCCCTCCTGC	TGGGCTCGCG	CTGGGAGATC	ATCCAGACAG	TCCCCGACGC	CACGCCAGgt	aggtcccttg
	C L L	L G S R	W E I	I Q T	V P D A	T P	
8840	caggagctgc	aggaggggggt	tgggcccccg	cagtgcattct	ccgatttcctc	cccattcccc	cacagAGTGT <sub>E C</sub>
	CCACATCAAG	AGATGGACAT	CGTCTTCTGT	ATTGACGGCT	CTGGAAGCAT	TGACCAAAT	GACTTTAACC
8910	P H Q	E M D I	V F L	I D G	S G S I	D Q N	D F N
8980	AGATGAAGGG	CTTTGTCCAA	GCTGTCTATGG	GCCAGTTGA	GGGCACCTGAC	ACCTGgtga	agactgggga
	Q M K G	F V Q	A V M	G Q F E	G T D	T L	
	INTRON 6 (1255 bp)						
10240	aacaatagta	acaggcactg	agccctgggc	cctcccact	ggcctttgca	gTTTGCACtG	ATGcAGTACT
10310	CAAACTCCT	GAAGATCCAC	TTCACTTCA	CCCAATCCG	GACCAGCCCG	AGCCAGCAGA	GCCTGGTGA
	S N L L	K I H	F T F	T Q F R	T S P	S Q Q	S L V D
10380	TCCCATCGTC	CAACTGAAG	GCCTGACGTT	CACGGCCACG	GGCATCCTGA	CAGTGGTgta	aagcaacccc
	P I V	Q L K	G L T F	T A T	G I L	T V	
	INTRON 7						
	gacccca						

Sequence Range: -11390 to 10387  
(SEQ ID NO:3)

**FIGURE 9**  
Protein Sequence: SEQ ID NO: 2

Translational stop codon for CDIlc

<u>TGAT</u> CCCTCT	TTGCCTGGA	CTTCTTCTCC	CGCGATTTTC	CCCACTTACT	TACCCTCACC	TGTCAGGCTG	-11321
ACGGGGAGGA	ACCACTGCAC	CACCGAGAGA	GGCTGGGATG	GGCCTGCTTC	CTGTCTTTGG	GAGAAAACGT	-11251
CTTGCTTGGG	AAGGGGCCTT	TGTCTTGTC	AGGTTC AAC	TGGA AACCT	TAGGACAGGG	TCCCTGCTGT	-11181
GTTCCCCAAA	AGGACTTGAC	TTCGAATTC	TACCTAGAAA	TACATGGACA	ATACCCCAG	GCCTCAGTCT	-11111
CCCTTCTCCC	ATGAGGCACG	AATGATCTTT	CTTTCCTTTC	CTTTTTTTTT	TTTTTCTTTT	CTTTTTTTTT	-11041
TTTTTTGAGA	CGGAGTCTCG	CTCTGTCACC	CAGGCTGGAG	TGCAATGGCG	TGATCTCGGC	TCGCTGCAAC	-10971
CTCCGCCTCC	CGGGTTCAAG	TAATTCTGCT	GTCTCAGCCT	CCTGCGTAGC	TGGGACTACA	GGCACACGCC	-10901
ACCTCGCCCC	GCCCCATCTT	TCTAAAATAC	AGTTCTGAAT	ATGCTGCTCA	TCCCCACCTG	TCTTCAACAG	-10831
CTCCCCATTA	CCCTCAGGAC	AATGTCTGAA	CTCTCCAGCT	TCGCGTGAGA	AGTCCCCTTC	CATCCCAGAG	-10761
GGTGGGCTTC	AGGGCGCACA	GCATGAGAGC	CTCTGTGCCC	CCATCACCCCT	CGTTTCCAGT	GAATTAGTGT	-10691
CATGTCAGCA	TCAGCTCAGG	GCTTCATCGT	GGGGCTCTCA	GTTCCGATTC	CCCAGGCTGA	ATTGGGAGTG	-10621
AGATGCCTGC	ATGCTGGGTT	CTGCACAGCT	GGCCTCCCGC	GGTTGGGTCA	ACATTGCTGG	CCTGGAAGGG	-10551
AGGAGCGCCC	TCTAGGGAGG	GACATGGCCC	CGGTGCGGCT	GCAGCTCACC	AGCCCCAGGG	GCAGAAGAGA	-10481
CCCAACCACT	TCCTATTTTT	TGAGGCTATG	AATATAGTAC	CTGAAAAAAT	GCCAAGCACT	AGATTATTTT	-10411
TTTAAAAAGC	GTACTTTAAA	TGTTTGTGTT	AATACACATT	AAAACATGCA	CAAAAAGATG	CATCTACCGC	-10341
TCTTGGGAAA	TATGTCAAAG	GGTCTAAAAA	TAAAAAAGCC	TTCTGTGGAT	ATGAGTCCTG	AAGGATGACA	-10271
CCCATGGGGT	CCCTTTACCA	CGGTGGACCC	TGGCCAGCAC	TGAGGCCTGG	GGCCAGGACA	AGAAGTTAAC	-10201
CAGAGTAGGG	TTGTGAATAT	CCCTCTCTTG	GAAGTAACCT	GACCTCTTAA	TCTGCTCACT	CCACTCTCAG	-10131

FIGURE 9 (continued)

GGCTGGTGCC	GATGGTAAGC	TGGTGGAGCT	GTCGGGTGGA	GGGGGCATAG	AATAGAGAAG	-10061 GGACAACCTC
CAGTGGCTAC	TTTTCACCT	GGAAAGGTCT	CTGGAGTGAC	CAATACTCAC	AAGCGTTTCC	-9991 TACAAGTCCT
AGGATGTGTT	GAAGGGCACA	CTGTCTGCAT	ATAGTGAGTG	ATTGAAGAAC	ATGTTGGGGT	-9921 CCCACATTGA
GAGCTGCTGC	CCACAATAAG	GTCATTCTTG	CTATTATGCC	ACCATCCTGG	CATAAAGTTC	-9851 ATCATGGTGC
TTGGCACTGA	GCTGGGGGCC	TCACAGGACA	AGCCATTCCCT	GACCTCGGAG	TGACGCCACT	-9781 GCAGCTATCA
CCAGCAAGGG	ACCCGGGCCG	TGTGGATGTT	TCAATTAGAA	AAACAGAAGG	GAGGCAGTTG	-9711 AGTGATTTGA
AGGGAAGATG	GAAAGTGGCC	CTTTACCTCC	AGCCAAAAAT	GTCTGTCCTA	TACATCAGCA	-9641 GAGGCTCCAA
AATCCCTGTG	GATTTTGAAG	CTTTTGAGTC	CCCAGGATGA	CTAATTATTA	TGCAGTTTCC	-9571 TCAGAAAGGG
AATCAGAAGA	TAAGGCTTTG	TAAGAATTCA	GCCCTAATGG	CTGGGCACAG	TGGCTCATGC	-9501 CTGTAATCCC
AGCACTTTGG	GAGGCCGAGG	CAGGAGGATT	GTTTGTGCTC	AGAAATTTGA	GACCACCCCTG	-9431 GGTAATATAT
TGAAACCTTG	TGTCTACAAA	AAAATTTAAA	AATTACCCAG	GCATGGTGGC	ATGTGCTTGT	-9361 AGTCCCAGCT
ACTTGGTAGG	CTGAAGCAGG	AGGATCACTT	GAGCCTGGGA	GGTTGAGGAT	ACAGTGAGCT	-9291 GTGATTTGGA
CCACCACACT	CCAGCCTGGG	CAACAGAGAA	AGATCATGTC	TCAGAAAAAA	AAAAAAAAT	-9221 TGACCCTAGA
GTGGTGT TTC	TCAAAATGTG	TTCCACGAAC	CACTGGTGGT	CAATGATGGT	CTTCTAAGTG	-9151 GAAGGTTTTA
GAGAAAAAGA	GCAAGAAACC	CATACATCTC	AAACATTTGA	AACTAGTGAT	TTGCACAGAA	-9081 ATAGTGTTGT
GGCCTTAATA	ATTGTGTGGC	ACACGGACTC	CAGGGACTAC	AGTGGGTTCT	TGTCTAAATT	-9011 CAGGCAACAA
GTTGTTATTT	TCTATTTTAT	TTTATTATTA	TTATTTTTTG	AGATAGTCTC	ACTTTGTCTC	-8941 CCGGGCTGGA
GTGTAGTGGC	ACGATCTCGG	CTCAACGCAA	CCTCTGTCTC	CTGGGTTCAG	GTGATGCCTC	-8871 TGCCTCAGCC
TCCCAAGTAG	CTGGGAGTAC	AGGGGCGTAC	CACCATGCCC	ATTTTTATTT	ATTTATTTTT	-8801 GAGACAGAGT

FIGURE 9 (continued)

CTCGCTCTGT	CACCCAGGCT	GGAGTGCAGT	GGCATGATCT	TGGCTCACTG	CAACCTCCGC	CTCCCAGGTT	-8731
CAAGTTCAAG	CGATTCTCCT	GCCTCAGCCT	CTGGAGTAGC	TGGGATTACA	GGCAGGCACC	ACCATTTCCTA	-8661
GCTAATTTTT	GTATTTTTAG	TATAGATGGG	GTTTCACCAT	GTGACTAGG	CTGGTCTCGA	ACTCCTGACC	-8591
TCATGATCCG	CCCTCCTCGG	CCTCCGAAAG	TGCTGGGATT	AGAGGTATGA	GCCACTGTAC	TTGGCCGACA	-8521
AGGTGTTATT	TTCTGATATT	CTTCCTTTGT	GTGTTATTGT	GTACATTTGT	TACATTTGCA	TTTTCAGGGT	-8451
TGGCTATTGT	GTTGCATTAG	ATCCCCGAAT	CACAAAATGG	ATCAATGGCT	CAAAAGCATG	GAAGTTGTGA	-8381
TTAAAACTA	ATCTAATTGC	TACAATTTAC	AATAATGTCA	TCAAAGTCAA	TATTGACTTT	TAAATATTGA	-8311
GCCCAGTGCA	CGTATAGTAT	AGACATGCAT	ACCGGAATAA	GTGATTGTGA	GCCAAAACCC	GAAAATATCT	-8241
AGAAGGTATT	ATACTCCCTG	ACAGGTAGGT	TGTATTGGTT	CTGACATGTA	TTTGTCCCTA	GTGTGCTGCC	-8171
CATTCTGAAA	CTTTATCAAA	CAGTCGCATG	AACCTCTGAA	AGCTTTTG TG	TTATTTTCTT	ATTTATTTAT	-8101
TTATTGAGAT	GGAGTCTTGC	TCTGTCGCCC	AGGCTGGAGT	GCAGTGGCAT	CATCTTGGCT	CACTGCAACC	-8031
TTTGCCCTCT	GGGTTCAAGT	GATTCTCCTG	CCTCAGCCTC	TTGAGTAGCT	GGGATTACAG	GCGCGCACCA	-7961
CCACGCCCAG	CTAATTTTTG	TATTTT TAGT	ATAGACGGGG	GTTTCACCAT	GTTGGTCAGG	CTGGTCTCGA	-7891
ACCCCTGACC	TCATGATCTG	CCTGCCTCAG	GTAAAGCAAT	AGAGATTCTT	AGAACAAC TG	CTACATGTAG	-7821
CTTTCCTATT	CAAAAGTGAT	TAGTGTGTC	ACCGAATACA	GAGGAGACAG	CAAAACCACA	GTGACATAAA	-7751
TCAAAGGTGC	TTTTTAAAGT	AGCAAAAGTA	GGTACAAGTC	ACATAATTTT	CAAGAAGCTT	GTAGAAATGG	-7681
CAGTAGAGTT	CATACCTGCT	ATTGAAAGGT	TGCTTTTGGC	TGCAAATAAT	AGAAAAAAC	AAAAGCATGT	-7611
AAGAGCAGAC	AGAAGACCTT	TACTCTGCAA	GAGGTTCAGG	TGCAGGTTAG	TGTTTAATGC	AGAGTCTCAG	-7541
CATTGACAGA	TTCTTTCTGA	TCTTCCAATT	GATCGTCCTT	GCGGGGGCGG	TTTAGTTCTT	TCCCACTGAC	-7471

FIGURE 9 (continued)

TAGGATTGGG	TCAAATTCCA	TCCCCTTGGT	TGCATGCAGT	GCTGAGAAGG	TGAGCATGTG	CTTTTCACAG	-7401
GCTTAATAAA	AAGAGGTAGC	TCCAGCCAGG	TGCAGTGACT	CATGCCTATA	ATCTCAGCAC	TTTGGGAGGC	-7331
AGAGGTGGGT	AGGTCACCTG	AGGTCAGGAG	ATTGAGAACC	AGCCTGACCA	ACATGGCAAA	ACTCTGTCTC	-7261
TACCGAAAAT	ACAAAAATTA	GCTGGGCATG	GTGGCAGGTG	CCTGTAATCC	CAGCTACTTG	GGAGGCTGAG	-7191
GCAGGAGAAT	CGCTTGAACC	TGAGAGGTGG	AAGTTACAGT	GAGCTGAGGT	CATGCCACTT	GCACTCCAGC	-7121
CTGGGGGACA	GAGTAGAACT	CTGTCTCAAC	AAAAAAAAAA	AAAAAAGAG	AAAAAAAAG	GAGGGTAGCT	-7051
CCACCAGCCA	GGAAGGTGGC	AGCGCTGGTG	GCTGTTGGAT	AGGCTACCTA	CAGTGTCTGG	CAAATACTAT	-6981
GCTTGAAGAC	TATGCTGTGA	GCAAGATTCC	TTTGTGAAGG	AACAGCTTGG	ACATTGTGTA	TGTCAGAGGT	-6911
ATACAGCAGA	ATAGCAGTGA	CTAACGCTTG	TGTGGGAGAG	CAAGCATGTC	ACCTCATACT	TGGAATAACT	-6841
CACTGCCATA	CAAAGTCTGA	ATCAGCTTTC	GTCTTTGTGC	AACACATGTA	TGTGGGAGCT	TTTCAGCTGC	-6771
TGAAACCTCT	AGTGACAGAA	AAGGAGGTTT	TGTTGTTCAT	TTGTAATTAA	TGTTAATCCT	ATGAGTGGTG	-6701
GGAGAGATAG	TGAGGTAGGA	GATCAGCAGG	ACCTGTTTTC	TGGTCACAAC	CCAGCTAATC	AGAGCATGAT	-6631
CTGGTCAAGA	TGGGATGCAC	TAAAAAACA	GCCCAAACCA	GCAGATGGCC	AGGAAAGCAA	ACTCTCATTA	-6561
CCCTCGCCAC	TTATTAGCAT	AAAGACACTC	CCACCGGTGC	CATGACAGTT	TACAAATGCC	ATGGAAACAC	-6491
ACCATAGCAA	CGGTCAGCAA	GTTACCTCAT	ATGGTTCTGG	AAACTCCCCA	CACCTTTTCC	AGATAGTTCT	-6421
GAATAACCCA	CCCCTTAATT	TGCATGTAAT	TAAAAGTCGG	TATAAGTACA	GTTAGCCAGC	AGCCCACTGG	-6351
CTGCTACTGT	GGGCTCACTG	CCTATGGGTT	GTCTTGCTCT	GCAAGGAACA	GCTACCTTGC	TGCCACTGCT	-6281
GCTTCAATAA	ACCTGCTTTC	TTCCACCACA	GGCTCGCTCT	TCAGTTCTTT	CCTGAGCAAA	GTTAAGAACC	-6211
CTCCCGGGCT	AAGCCCCAAT	TTTGGAGCTT	GCCTGCCCTG	CATCAGTAGA	ATGGGCTAAC	TACTTACGGT	-6141

FIGURE 9 (continued)

GCACTCAGGC	TAAAGAGGCT	GATGCTTGCA	GGGCAGTATT	CACAGAGCAC	ACGGTAGTTC	-6071 ACGGGATGCC
TCTCACCCCTT	GACTCAGTGC	TTAAGAAAGG	AGGGAAAATG	GTGAACATGA	TCAAATCATG	-6001 GCCATTGCCT
ATTCATCTTT	TCAGTGTTGT	ATGGAGGAAT	AGGCAAGTAG	GAGATTGCTT	TTCACATTAA	-5931 TGTCAAAGAG
AAAGATAGTT	ACTTGGAAC	TAAAAAATT	AATTGTGATA	AAATATACAT	AACATAAAAT	-5861 TTACCATCTT
AACCATTTTT	AAGTATAGCC	AATCTCAAGA	GCTCTTTCTA	TCTTGTAATA	CTGAAACCCT	-5791 ATACCCATTA
AACAACCCC	AATTCTCCCC	TTTCCCTAAC	TCCTGGCAAC	CACAATTCTT	TCTGTCTCTA	-5721 TGAATTTGAC
TGCTTTGGCA	TGTCATAGAA	ATAGACTCAT	ACAGCATTTG	TCTTTTGGCG	ACTGGCATAT	-5651 TTTGCTTAGC
ATAATGTCCT	CAAGGTTTAC	CCATGTGGTA	GCATGTGTCA	GAATTCCTCT	CCTTTTGAAG	-5581 GCTGAATAAT
ATTCCATTGT	GTGTATATAC	CACGTTTTGT	TTATCCATTT	GCCCATCAAT	GGGCATTTGG	-5511 GTTGCTTTTT
TTGCCTCTCA	TGAATGATGA	ATATGGGCGC	ACAAATATCT	CTTCAAGACC	ATGCTTTCAA	-5441 TTCTCTGGG
TATACACCCA	GAAGTGGAAT	TGCTGAATCA	TATGGTAATT	TTTTTTTTTT	TTTGAGACAG	-5371 AATCTTGCTC
TGTTGCCCAG	GCTGGAGTGC	AGTGGCACAA	TCAGAGCTCA	CTGCAGCCTT	GGTCTTCTGG	-5301 GCTCAAGCGA
TCCTCTTGCT	TCAGCCTTCC	GAGCTTCTGG	GAATAAAGGT	GTGTGCCATC	ATGCCTGGCT	-5231 AATGTTTTAA
AAACGTTGCC	AGGCATGGTC	GCTCGTGCTT	GTAATCCTAG	CACCTTGGGA	AGCTGAGGCA	-5161 GGTGGATCCC
CTGAGGTCAG	GAGTTTGAGA	CCAGCCTTGC	CAACATGGTG	AAATCCCGCC	TGTACTAAAA	-5091 ATACAAAAAT
TAGCTGGGTG	TGGTGGCATG	TGCCTGTAGT	TCCAGCTACA	GGCAGGAGAA	TTGCTGGAAC	-5021 CTGGGTGGCA
GAGGCTGCAG	TGAGCCGAGA	TTGCACCACT	GCACTCCAGC	CTGAGTGACA	GAGTGAGACT	-4951 CTGTCTCAAA
AAAAAAAAAA	ATTTTAGAGA	TGGTGTCTCA	CTGTGTTGCC	CAGGCTGGTC	TTGAACTCCT	-4881 GCCCTAAAGT
GATCCTCCTG	CTTCCGCCTC	CCAAAGTGCT	GGGATTACAG	GCATTAGCCA	CCATGCCTGG	-4811 CCTAGCTAAA



FIGURE 9 (continued)

TTGTCTTTAA	TGTCGCATGT	CTGCAAAAAA	CACATCTATA	AAGCTAGAAA	AGTTGAGCAT	CCAACTTTTT	-4741
ATGATTTAAC	TCTCATGACC	TGGCAATTTT	TCTAGCAAGG	AGCCTGGGCT	GGTGGTTTTA	GGAGAACTGA	-4671
GTGAAAAAAA	GAAATACATT	AACTAGATTG	GATGCAAAGT	GCCTGCTGGT	CATGGGTGTT	TTCTGCTGGC	-4601
CCCTGTTTAT	CTGTGCCTGT	TAGCCCACCC	ATGGGTGAGT	GGGGCAAAGT	GGCCAACTG	ATTCTTAAGA	-4531
GAGGCATACA	TGCAGAATCC	AAGTTAGTCA	TGATTTCTGT	TCTAGTCTGA	GTGAATGTGT	GTCCAGAATA	-4461
TTTTATAAAC	TTTATCAGCT	CAGAGGGGAA	AACCTGTCTC	CATACTACGT	GGTTTATACA	AAGCTGTCAG	-4391
GAATTCAGCA	TGATGAAGAA	ATGCACAAAA	CAAGTGTGAA	CAGATAAGTA	AAAGGATCTA	CTGAAAATCT	-4321
TCAGGGTAGT	ATATTGTGTG	ACAGGACCAA	GAATTTGAAG	TCAACATCTG	TATTTGTGCC	CTCTGGACAA	-4251
AGGTATTATC	CCTGATGATA	TAAAAATTAA	TTTTGGGCTG	GGTGTGGTGG	CTCATGCCTG	TAATCCCAGC	-4181
ACTTTGGGAG	GCTGAGGAGG	GTGAATCGAC	TGACGTCAGG	AGTTGGAGGC	CAGCCTGTAT	CGACTAATAA	-4111
TACAAAAAAA	TTAGCTGGAC	ATGGTGGCGT	GCACCTGTAA	TCCCAGCTAC	TCAGGAGGCT	GAGGTGGGAG	-4041
AATTGCTTGA	ACTCGGGAGG	CCCAGGTTGC	AGTGAGTCGC	ACCACTGCAC	TCTAGCCTGG	GCGACAGAGT	-3971
GAGACWCCGT	CTCAAAATAA	ACAAAATTAA	TTTCGAGGCC	AGGTGCAGTG	GCTCCAGGTG	CGGTGGTTCA	-3901
TACCTGTAAT	TCCAGTGCTT	TAGGAGGCCA	GAGGATTGCT	TGAACCCAAC	AGTTCGAGAT	CAACCTGGGC	-3831
AACATCAGTG	AGACTCCATC	TGTAGAAAAC	AATCAAACAG	ACAAACAACA	ACAACAAAAA	AACCAGAGGT	-3761
GGGAGGATCA	CTTGAGCCCA	GGAGTCCGAG	GCTGCAGTGA	GCTATGGTCA	CGCCACTGCC	CTCTAGCTTG	-3691
GGCAACAGTG	CCAGACTCTG	TCCTTAACAA	CAACAACAAC	AAAAATTAAT	TCTACTTTAA	CTGTCAGTTT	-3621
CATGATATCC	TTCTATTAAG	AAAAACCTTT	TCTATCTGAT	GAACATTGG	CTAGGTTTTC	TTTCTCTCTG	-3551
CTTTTGACTA	ATGCATTTAA	TTACTTTCAT	TTGCAAACTC	TATCCTTCTC	ATCAACTTTG	TATTTTAGAT	-3481

FIGURE 9 (continued)

GTGTCTATTG	ACAGCCTGGC	TCCCTCAGC	GATCATTATG	ATGATCAAAG	TAGATGAATA	GGTAAAATTC	-3411
AATGCAAATA	TTCCAGGGCA	TCTAAATCCA	TACCCCAAAT	GGGAAAAGGG	GAGAATTGGA	AGCCAGCAAT	-3341
TTGAACACAT	TACTATGGAT	GTATTTTCT	CATGCGGGG	AAAAAGTGAT	TTGGAGAGAG	AGAATTATGA	-3271
ATGCATGTGA	AGAATAAAGC	CAAATTCCT	GGGAGGAGGG	GAAGACCAGG	AGAAACAAAA	CCAAATCCTG	-3201
GCTGTGGCCT	CTAAGGCATG	GGGACCTGGA	GTTATGCTCT	CCAGGCAGAC	ACAGCTCATT	CTGGAGAAAG	-3131
GCTGCAAAAA	TATTCTCCTT	CACATTGATT	TGAAAACAAT	TATTAAATTC	TTGTTTTCTT	ATTTATCTAA	-3061
GTGTAACTTT	TTAAACTTA	CTGAGAGAAG	ACGGGCACGG	TGGCTCACTG	CTGTAATCCA	GTACTTTGGG	-2991
AAGTCAAGGC	AGGTGGATCA	CCTATGGTCA	GGAGTTCGAG	ACCAGCCTGG	CCAATATGGC	AAAACCCCGT	-2921
CTCTACTAAA	AATACAAAA	TTATCAGGTG	TGGTGGTGTG	TGCCTGTAAT	CCCAGCTACT	CGGGAGGCTC	-2851
AGACAGGAGA	ATCACTTGAA	CCTGGGAGGC	AGAGGTTGCA	ATGAGCTGAG	ATTGCACCAC	TGCACTCCAG	-2781
CCTGGGCGAC	AGAGCAGGAC	TCCATCTCAA	AATAAAAATT	AATTGATTAA	TTAATTAAAA	ATTTACTGAG	-2711
AGCTGGTGGT	TCCTTTAAGG	GTGGAGCCGC	CATCAAGTCC	CCAGAGGATG	CCCTGAATTT	GGGGGCATCA	-2641
CCTTCAGCTG	CTGTGGACTC	TGAGCCTTGG	CAGCTCCAGC	TCCAGGCCTG	GGAGAAAGAT	GATTTCTTGG	-2571
CAGCGTGCAG	TGATTGTGAG	CATTTGACTA	CCTTACTGCA	TTTGCCCTT	ATCAKTGCTC	TCCAAACATG	-2501
AGTGGAAC	AAAAAATTTT	GCTGAGACAA	GCGATAATAC	GAGTTAGGGA	AAGTTGGAGA	ATTTTATAGT	-2431
TGCTGATATC	AGCAAATCGT	GAGTTTCAAG	CACTAACTTA	CAGAAGGAAG	TCCAAAATTA	AAGGGGATAT	-2361
AGAAATGTGT	AAAAGATGAG	GTGTGGTGAA	GATGGAGAAA	ATGAAGAGCT	CTTTAAATTT	CTGAATTATG	-2291
AAGAATCACC	AACAAATTAT	TTTGTGGTTC	CAAATACAGG	GAGAAGTTCA	CAGATCCACA	GAAGTATGTA	-2221
CAGGGTGCGG	CCAGCCACAA	ACCTTTCAGC	ACAAGAGGGA	GAAGGCTGCC	GCTCCACTTT	GCCTGGGCAG	-2151

FIGURE 9 (continued)

TCTTTGTAAG	GCAGTAGATA	AGTCAGCCTC	GAAGTTAGCA	ATCACAGCCC	TCGGCTCGGT	-2081 TTCCTGCAAG
GGCATCGTTA	ATGCATCACA	ATTAATTTCT	TCTGTCCATT	AAATGTCAGC	TCTCAAGTAA	-2011 ATTGATGTAA
AATTTTGTGA	TAGAAACTA	TTTCATATTA	TTTGCACTTG	ATGTTTAATT	ACATTTTAAA	-1941 TGTTTTGTTT
GTTTCATTTT	GTTTTGTTTT	TGAGACAGAG	TCTTGCTCTG	TTGCCCACGC	TGGAATGCAG	-1871 TGGTGTGATC
TTGACTCACT	GCAACCTCTG	CCTCCTGGGT	TTAAGCGATT	CTCCTGCCTC	AGCTTCCTGA	-1801 GTAGCTGGGA
TTACAGGCGT	GCACCACCAT	GCCTGGCTAA	TCTTTGTATT	TTTAGTAGAG	ATGGGGTTTC	-1731 ACCATGTTGG
CCAGGCTGGT	CCCGAACTCC	TGACCTCAAG	CTATACACYT	GCCTCAGCCT	CCCAAAGTGC	-1661 TGGAATTACA
GACATAAGCC	ACTGTGCCCA	GCCAAATGTT	TTAAATAATT	GTCACATATA	TATACAAAAT	-1591 AATTTATGTT
ATAGGTAGGG	ATCTTGTTAT	ATTTTAACCT	TCAAAGTATA	TTCCTAAGCT	TTTTATTTAT	-1521 TTTTTATTTT
TTATTTATTG	AGACAGTCTT	GCTCTGTCGC	CCAGGCTGGA	GTGCAGTGGC	GCAATCTCGA	-1451 CTCACTGCAA
ACTCTACCTC	CTGGGTTCAA	GCGATTCTCC	TGCCTCAGCC	TCCTGAGTAG	CTGGGATTAC	-1381 AGGTGCGCAC
CACCATGCCC	AGCTAATTTT	TGTATTTTTA	GTAGAGACGG	GGTTTCACCA	TATTGGCCAG	-1311 AGCTGGTCTC
AAACTCCTGA	CCTCAGGTGA	TCCATCCACC	TCAGCCTCTC	AAAGTGCTGG	GATTATAGGT	-1241 GTGAGCCACT
GCGCCTGGCC	TATTCCTAGC	CTTTTATATA	TAGACCTTTT	TCTTTTTCAC	ATTTTAAAGG	-1171 AACTTTTATG
TTTAATCATG	GAATATTTCA	AACATACAGA	AAAATCACAG	AAAATAAATA	ACAACCACTC	-1101 ATTTATCTTC
TCCCCAACCC	CATGTAATAA	ATATTAAAAT	ATTGTGTTAA	ATGCTAAATT	TAACACATGC	-1031 TAAAGGTTC
TGGCTGGATG	TGGTGGCTCA	CGCCTGTAAT	CCCAGTACTT	TGGGAGGAGG	AGGTGGGAGG	-961 ATTGCTTGAG
TCCAGGAGCT	CGAGACCAGC	ATGGGCAACA	TAGTGCGATC	TCGTCTCTAC	AAAAAACAAA	-891 AAAATTAGCT
GGGCATGGTG	GTGTGCATCA	GTAATCCCAG	TGACTGGGAG	GCTGAGGTGG	GAGAATTGCT	-821 TGAGTCTGGG

FIGURE 9 (continued)

AATTTGAGGC	TGCAGTGAGC	CCTGATCATG	CCACTGCATT	CCAGCATGGG	CGACATAGCA	AAACTTGTCA	-751
AAAAAAAAAA	AAGTTTCCTC	TCTGCCCCAC	CATAGACAAC	CACTCTTCTG	ATTTCTATCT	TCGTAGAT68	-681
ATTTTGCCCA	TTCTCTTGTA	TATGAAAGAA	AP1 ACCAGACATT	AGGCATTCTG	GTGTCTGGTT	TCTTTCACCT	-611
AAGATAAAAT	TGAGTTAACC	TGTATTGTTG	TACAGAACTG	CAGTTTGTTC	TTTGTTATTT	ATTGTAAAGA	-541
CAGGGTCTGG	CTATGTTGCC	TAGGCTGGTC	TCGAACTGTT	GGCCTCAAGC	AATCCACCTG	CCAAGCTCTG	-471
GGACCACAGG	CATGAGCCAT	GGCATCTGAT	CKGTAGTTTG	ATCTTATTTT	TTGCTGAGTA	GTAGCCCATG	-401
AP1 GCATGACTTT	ATTATTTTGG	GTGTCCATTC	TCCTCTGGAG	GGGCTCTGCT	TTTTGAAACC	ACACCCTGGC	-331
Ets CTAGCTCCCC	TTCTCCCTGC	CTCTCTGCAG	GCTCACATCC	ACATGCCAAG	ACCTCTGCAG	CCATTCTGCT	-261
Ets TCCTGTCCCT	CCACTCCTGT	GGGACCTCAG	AGAGCTACGG	GGCTCCCTGG	GTACCAACTG	GCTCCTGAGG	-191
Sp1/Sp3 CCTGGGGGAG	GGTGGTCTTC	TGGGAGAAGG	AAGCCAGGTC	CCTGCAGGTT	Sp1/Sp3 GTGGAGGGGG	ACAGAATGAG	-121
GGTTTTTCCC	CAGGATGTTG	Sp1/Sp3 TTGGCCCCCTG	CCCCCACTTC	TGTTCCATAA	TTAACCACGC	Ets CCCTCCTACC	-51
CACTGTGCCC	CTCTTCCTGC	Sp1/Sp3 TGTGTGGAGG	CCCTGAATCA	TTATTTTAAC	+1 TACCCCTGG	20 GAGGGTGAGC	
Ets ACCTTCTGTG	CTCTGTCCCC	Ets AACCTTCCAC	TTCCCCTCAA	CGCGCTGCTC	AGGGATGACC	90 TTGGCACTG	
TCCTTTTCT	GAGTGGTAAG	TGGGGCCAGG	GTGCTGGGGA	GAAGCTTGGA	GGAGTTCTGA	160 GGGGACTCCA	
V L L L	S V						
TCTGGGAGGG	CAGGCTGGGG	GCTGGTGGTC	GGCTCCAACC	ACTCTTATGA	GGAGCTGAGG	230 CAGGGGAGTG	
CTTCATGTGC	GAGTGGCCCG	GAGTCAGTAG	AGTGTGACCT	GAATGAAGAG	GGGCTCAGGG	300 GCTGTGCTCA	
GGTGGCGACT	AAGCTACCTC	TCCAGCTGGC	TATGTTGTCC	CAGGCTTCCC	TGCTCCCACT	370 CATGGAGTCC	
CTGGTGTGGG	TGACAGAGGT	CTCCCCAGCC	TCCCCCGGA	GTGGAAGGCC	ACAGAAGCCA	440 CCAGGGAGGG	
GGAAAGGTTG	GACATCACCT	CCCTGGGCCT	NNNNNTTCCC	CCAAGTCCTG	ACTGCACGTA	510 GGGAAGAGGC	

FIGURE 9 (continued)

CCCCTGCTGA	AAACTGCATC	AGAGTCACAT	TCACGTGCCA	TCAAAAATCA	GGCTTGGCTG	GGTGCGGTGG	580
CTCATGCTTA	TAATCCCAGC	ACTTTGGGAG	GCCGAGATGG	GCGTATCCCC	TGAGGTCAGC	AGTTTGTGAC	650
CAGCCTGGCC	AACATGGTGA	AACCCCATCT	TTACCAAAAA	TATAAAAATT	AGCCGGGCAT	GGTGGCGTGC	720
ACTTGTAATC	CCAGCTACTT	GGGAAGCTGA	GGCAAGAGAA	TCGCTTGAAC	CCAGGAGACG	GAAGTTGCAG	790
TGAGCTGAGA	TCGTGCCGTT	GCACTCCAGC	CTCAGCAACA	GAGCGAGACT	CCATCTCAAA	AAAAAAAAAA	860
AAAAAAGAA	AAAAAAGAAA	AAGAGGCTGG	GAGGTCCTAG	GGATTGGGGC	TTCTTTAACT	CCCAGCCTCC	930
CCGCCCACCA	AATATTCCTC	AGTCTTGCT L A	TCTTATCATG S Y H G	GATTCAACT F N L	GGATGTGGAG D V E	GAGCCTACCA E P T	1000
TCTTCCAGGA I F Q E	GGATGCAGGC D A G	GGCTTTGGGC G F G	AGAGCTGGT Q S V V	GCAGTTGGT Q F G	GGATCTCGGT G S R	AGGCCCCACT	1070
CACCCTCCTT	CCCCAACCTC	CACTACATCA	AGTCCTGTGG	ATGGGTACAC	GTGGGTACC	CGAGGGAGGT	1140
GTCCTGGAGG	AAGGCCAGCA	GGGGTGAGAA	GTCTTCCCTT	GGCTCCTTGG	AGGCCCTGAC	ATCAGCACCT	1210
ATTATTCTCA	ATCCCAGGAA	AGGCCACAAA	ACTCTAGACA	AGACCCTACC	TTACCTCGGG	AGGGAAGCCT	1280
TGAACCTGCC	TCCCAGGCAG	GGCCCACTTC	TTGGGGCCAG	TATGGTCACA	CAGGGCCAC	ACTCATTAAC	1350
TTTGAGTTT	AATGTTCTGC	CCTTGACCTC	TTGAAATTCC	TGATTATTTT	TATTTTATT	TTTACTCCAG	1420
CTCTGTTACC	CAGGCTGGAG	TGCAGTGGTG	CAATCACAGC	TTACTGCAGC	CTCAAACCTCT	CGGGCACAAG	1490
TGATCCTCTC	ACCTCAGCCT	CCTGAATAGC	TGGGACCACA	GGTGCATGCC	ATCATGCCTG	TTTTTTGTTT	1560
TGTTTTGTTT	TACTTTTAC	AGAGATGGAG	TCTTGCTATG	TTGTCCAGAC	TGGCTGAACT	CCTGGGCTCA	1630
AGCAATCCTC	CTGCCTTGGC	CTCCCAAAGT	GCTGGGATTA	CAGGTGTGAG	CCACCCTGTC	TTGCCAATTC	1700
TTAAAAATTT	TATCTGTGCA	TTGTGTTTT	GCAAGTAAAG	AATGATGGCA	GGGCTGGGCA	CCATGGCTCA	1770
CGCCTATAAT	CCCAACSCCT	TGGGAGGCTG	AGGCGGGCAG	ATCATCTGAG	GCCAGGAGTT	TGAGACCAGT	1840

FIGURE 9 (continued)

TTGGCCAACA	CAGCAAAACC	CCATCTCTAC	TAAAAATGCA	AAAAAAATTA	GCCGGGCATG	1910 GTGGCAGGCA
TCTGTAATCC	CAGCTACTTG	GGAGGCTGAG	GCAGGAGAAT	CGCTTGAACC	TGGGAGGTGG	1980 AGGTTGCAGT
GAGCCGAGAT	CGTGCCACTT	TACTCCAGCC	TAGGTGACAG	AGTGAGACTC	CGTCAAAAAA	2050 AAAAAGTCAT
GGGAGAAGGG	AGATGTACTG	GGGGTTTGGA	GCCTTAGCTC	AGCAGCAGCC	CCACCTCCCA	2120 CCGCCTCCTG
AAGGGTGGTG	AAGGGGTATC	AGCTGCTGGC	TCCCCACCC	ATGTGGGAGC	AATGACCGCT	2190 GCTACCTTCC
GCCCCTGGA	TGAGCTGGGT	AAAGTCAGTT	AGGGGCGCTC	ACTCTGGGAG	TACCCCGAGG	2260 GAGTGGGACA
CTACATAGCA	AATAAAAAAC	GTCAGGACAG	GTTGAGGAAA	GAGAGCAGAA	GAAAGGTAAG	2330 AGCCCCCAA
CCCCAAGAGA	CCCCACAGTT	TTATTTCAAA	TTGGGACCCA	CAAATTATGA	ACCTGCCCCC	2400 ACTTCCAGGA
GCTCACATTC	TCCTGTCCCA	GAGAGTTCAA	GTCACAATGT	GACACAGGTG	TCACCAAGGT	2470 CTGGGGGGCG
CAGGCAGGGA	GAGAGCAGAC	CCAGGAGGGT	TCCATGGAGG	AAGTGGTGCT	GGCAGTGAGC	2540 CCCAGTGGAC
AGGAAGGCTC	AGTTGGTCAC	GAGGAGCTAT	AAGAGGTCAC	CGAGCTCCAA	CCGCGCACCC	2610 CTCTCCCTTC
CTCATGTGAC	TGGCAGTCTG	GGGGGATGGA	AGCAAGCACC	AGGCACCAGG	CTTTTGTTTT	2680 TCTTTATTTG
GAAATGTGGT	CAACTGAGGT	GCACAAATCT	GAAAGACCCA	ATCTGATAAA	GGATACACAT	2750 GTGCGTGCCT
GGGTGAGCCC	CACCTAGGTC	AGCTGCTCCA	GTGTCAAATC	CCACAGGCAC	AGGGCTGCCG	2820 TGGACCCCTT
CTCATCACCC	AACATCCCCA	GAGAACCCCT	GGTCAGACTT	CTGTCACCAT	CAGTTTTTTG	2890 GGCCACATTT
TAAAAAAGA	ATACATTGGC	TGAGTGCAGT	GGCTTATGCC	TATAATCCTA	GAACCTTGGG	2960 AGGCTGAGGC
GGGTGGATCA	CCTAAGGTCA	AGAGTTCAAG	ACCAGCCTGA	CCAATATGGT	GAAACCCTGT	3030 CTCTACTAAA
AAATACAAAA	ATTAGCCTGG	CGTGATGGCA	GGTGCCTGTA	ATCCCAGCTA	GCTGGGTGAC	3100 TGAAATAGGA
GATTTGCTTG	AACCTGGGAG	GTGGAGGTTG	CAGTGAGCTG	AGATCACGCC	ATTGCACTCC	3170 AACCTGGGTG

FIGURE 9 (continued)

ACAGAGTGAA	ACTCTGTCTC	AAAAAACATA	TGGGTTGATG	GGTTACACTA	AAGTTTGTGCT	CATCGTTTGT	3240
ATCAGCAGGT	TCCAAACTGC	TACCTCTCTA	GCCAATGCTC	AGATTTTCTT	CACAAAGCCT	TAGGCATCCC	3310
CTGAATCATG	ATGCACAGGG	ATTGTAGCTT	TCTGTAAAGG	AGCGGCACCT	AGAAGGAACC	CTCACATGGC	3380
CATTTAATGA	AGCCTTGCTT	GGCGCATTAA	AATACACCAG	ATATCTGTCTG	CTTTTCTCAC	AGACAGGAGA	3450
TTGTGGGTAG	TGAGAAAACA	TTTCCAAAT	TAAAAAACTT	TCCCACTCAG	GGAGTTTTCG	AAATAAACCC	3520
TTGACTCTAC	ATAACTATAG	ATATAGTTAT	GGATCCTAGT	ACACTGCTTT	ACATTGGCCA	ATTGAAATTG	3590
CTTATACAAT	ATTTAAATTG	GTCCAATGAA	TTACAGAATC	AACTATTGT	TTTGAAAGCA	CATGTCTTCA	3660
GGAAATTGTT	CCAATTAAC	TGAGATGATC	TTATTTCCTG	GGTGGTCAA	AATAATGGCA	ACTCAGAAAC	3730
GCAATGTGCT	TACCCATGAT	TGGGAAATGC	CATTTTGGTC	TTTAAATAGG	TCTTTTTTTT	TTTTTTTTTT	3800
TTTTTTTTTT	GGTGAATGTT	AAAAGAAAT	TTCTAAACAT	AAATACACAC	ATACGTACTT	ATGCACACTC	3870
AAAACCAAAT	AAACCCAGC	ATGGCCCCTG	GGCATCTGTG	AGTTACACTT	GGGCCCTGAT	TTCTGAATAT	3940
TCTGCCAAGT	GGCAAATGCC	AGGAATTTCC	CCCACAGAGT	CTCGCTTCCC	CATGGAGGGA	CACTTCCTCA	4010
CCCCCAAGTG	CCCGCTGCTC	CCACCCCTCC	TGTGGCTGCA	GTGACATGGC	CATGGTTGTG	TCTCCAGACT	4080
<del>CTGTGTGGA</del>	<del>GCACCCCTGS</del>	<del>AGGTGCTGTC</del>	<del>GGCCAAACAG</del>	<del>ACGGGACGGC</del>	<del>TGTATGACTG</del>	<del>CGCAGCTGCC</del>	4150
V V G	A P L E	V V A	A N Q	T G R	L Y D C	A A A	
<del>ACGGGATGT</del>	<del>GCAGCCCAT</del>	<del>CCCGCTGCAQ</del>	<del>AGTGAGTGAC</del>	<del>CACCTGGGAA</del>	<del>TTGGGCCCT</del>	<del>CAACCCTCCT</del>	4220
T G M C	Q P I	P L H I					
GGACCCAAT	GTGCCCCCGC	TTAGCTTCCA	Intron 3 GTCCAGACCT	TCCCCGCAAA	TGAGTGTGTG	CTGTGAGTGA	4290
GACCCCGCGT	GTCTGCCCTT	<del>GCAGTCCGTC</del>	<del>CTGAGCCCT</del>	<del>GAACATCTCC</del>	<del>TTGGGCTCA</del>	<del>CCCTGCAGC</del>	4360
		R	P E A V	N M S	L G L	T L A A	
<del>CTCCACCAAC</del>	<del>GGCTCCCGGC</del>	<del>TCTGTGTGAG</del>	<del>TGAGTGTCTT</del>	<del>GGGCCACGGG</del>	<del>GGGGTGGGGT</del>	<del>GGGGCGGGGG</del>	4430
S T N	G S R L	L					
GTGTTGTTGG	GGAGGAGGCT	GGGGCTGGGA	GTGAAGGAGG	AGGGGCTGCT	AGGGACTCCT	GGCTCACAGG	4500
			Intron 4				

FIGURE 9 (continued)

CTTCTGCCTC	CAGGCTGTG A C	GCCGACCT G P T L	GCACAGATC H R V	TGTGGGAGA C G E	ACTCATATC N S Y S	AAAGGTTCC K G S	4570
TGCTCTCTC C L L	TGGGTCGCG L G S R	CTGGGAGATC W E I	ATCAGACAG I Q T	TCCTGACGC V P D A	CACGTCAGGT T P E	AGGTCCTGG Intron 5	4640 (1700) (4267)
CAGGCCATGG	TTCCCTGTGG	AGCACATGCT	GGCACTGAGG	GTGAGCAGGC	GTGAGGCCTG	TGTCTGGGQC	4710
CCTGTGCCCT	CCCTGGAGGG	CCGAGTGTGG	CTAGGAGAGA	AGCCAGGAGA	AGAGGGTGGC	TCAGGCAGGA	4780
GCCCTGCTGC	TCCAGGGTAG	AAGTTCTTTG	CAGGGTTTTT	CTTTATATTT	TTTTCTTTTT	AAGACAGGGT	4850
CCCTGCCAGG	CACAGTGGCT	CAGGCCTGTA	ATTCCAGCAT	TTTAGGAGGC	TGAGGTGGGC	GGGATCACCT	4920
GAGGTCAGGA	GTTGAGACC	AGCCTGGCCA	ATGTGGTGAA	ACCCCTCTAC	TAAAAATACA	AAACAAACAA	4990
AAACAAATA	GCAGGATGTG	GTGGTGTGCG	CCTGTAATCC	CAGCCACTCG	GGTAGGCAGA	GACAGAAAGAA	5060
TCGCTTGAAC	CCAGGAGGCG	GAGGTTGCAG	TGAGCTGAGA	TTGTGCCATT	GCACTCCAGC	CTGGGTGACA	5130
AGAGCAAAAC	TCCATCTCAA	AAAAAAAAAA	AAAACAAAAA	ACAGAGTTTC	TGTCAGGCTG	CATGCACCAC	5200
CACACCCTGC	TAATTTTTTT	GAGACAGAGT	CTTGCTCTGT	CGCCAGGCT	GGAGTGCAGT	GGTGCAATCA	5270
TAGCTCACTG	CAGCCTCGAA	CTCCTGGGCT	CAAGTGATCC	TCCTCCCTTA	GCCTACTGAG	TAGTTGGGAC	5340
TGCAGGTACA	TGCATCACAC	CTGGCTAATT	AAAAAAAATG	TTTTTGTAGA	AATGGGGGTC	TTGCTATGTT	5410
ACCCAGCCTG	GTCTTGAAC	CCTGGGCTCA	AGTAATCCTC	TGCCACAGCC	TCTCAAAGTG	TTGGGATGAC	5480
AGGCATGAGT	CCTTGTGCCT	GGCCTGAGGG	ATGAAAGTTC	TGATGGAGGC	AGAGAGGAGC	CCCACTGTGC	5550
GGGCTGTAGA	GGGCACAGCA	TCTTCCAGTT	GCCAACAGGT	GCATGGCCAC	TTCTTGAGTT	TCAGAGGAAG	5620
GACCTTAGTG	TGGTAAAGAA	CGTGGTGAGG	AAGATAAATC	CATGAGGGAG	GTGTTTCTTC	TGGATGGTTC	5690
ACTGCTGAGC	TTCCAGGATT	CCCCAACTA	ACTTTCCTCT	CGAAGAGGAG	CAAATGACAG	GGCTGCGGAA	5760
AATGCGATGT	GCAATTTTGT	CAGTGCCCAT	GTCTTCCACA	GAGAACAGGG	CCTGGGGACA	CCACCATGAC	5830



FIGURE 9 (continued)

ATCTCTCTGA	GGGTGGTCT	GCATCATGGT	GGTCCCAAG	TTGTGTTTCC	ATGGGCACCA	GGCTTCATTC	5900
CCTTGAAGCT	TCATTCCCTC	AAAGCCATTC	AGTTTCCTCA	TTGGTAAAAT	AGAGCTCAAT	AATCAGGGGG	5970
TTATGAAGGT	GAAAGGGATT	GAGGTGCATA	AAGCACTTGG	AACCCTGCCT	GGCACATAGT	ATGTGATAGC	6040
CCCTCTGACC	CATCTTCCAG	CTGGGGACTG	CATGCTGGGA	CTGGGAGGAA	GATACAGGCA	AACTGTCTCA	6110
TCTGCCGTGT	GAGAGGGAAT	GCCAGGGGCC	GCTCAGGGTG	CTGACCGAGG	GTGGGGCTTC	AGACCAGAGA	6180
GGCCATGATG	ACAGGCATGC	TGGGCCTTTA	GACAAAGGTG	GAGCAGCAGC	AGAAACATTA	CCAGAGCAAA	6250
TGGTGAGGGT	GGAGTCTATG	GAGGGGACCA	AGGGAAGGGG	GAAGGGACAT	CCAGGGTTCT	TGGGGGGACC	6320
GTGCCCAGCC	TGAGATGTCT	GTGAAGCTAG	GTTAGGGAGG	TGGCACTTAA	AAACAAGGGG	TAAATGCTTT	6390
CTCACAGCCA	TCCGTGGAAC	TCATGAGGTG	GGATGCCTGA	TGCAAATGGG	ACTGGAGCAC	AAAACGGTG	6460
CAGGCAAGGG	GGGTGTGGGT	CCAAGTAGAA	GGGACCAGGG	TCCACTGAGG	ATCACCTGTG	TGCCAAGCAG	6530
TGCTGAATAC	CTGGTATGAA	TCACCTTATT	GCATCCTCAC	AACATCCTGG	GTGGTGGGCA	GGCCCATCT	6600
CATTTTACAG	ATATGAAAAC	CAAGGTTTCA	ATAGATGAGT	TCCATCGATA	GCAAGAGGCA	GAGCCCAGAG	6670
CTTGAGCCAT	CCTTGCCTGA	TTGGTGGGGT	CCTTTTTCAT	AAGGATAAGT	CCAGGCTTCT	GCTAGTGGGA	6740
GACCAGGGGA	TACAATAAAA	AGACCAAGAA	ACAGAAGAGA	CATTGTGAGA	GGATTTGCCA	CAGACCTGGC	6810
CTGAGAGAGG	ATGAGAGGGT	GGTTTCTTGA	CGCAGCTGAA	AAAACAGGCA	CCACTGCAAG	ATGTTGGCTG	6880
CCCAGATGTG	GGCAAAAAC	GGGGAGCTCC	TGGGGGGATC	TGCAGCCTGC	CCCATGGATG	TCAAGATTTG	6950
CTGGTGATTG	AAGAAGCAGG	AAGGAAGTGA	CCTTCTGTTT	CTCCCCAGCA	CCCTTGAAGC	ACCAGTGTTT	7020
GAGCAAGTGG	GGTAGGGGAG	AGGAAAGAGG	AAAAGGCATT	TTTTTTTTCT	GCAGTGGTGG	GCAGGGGGCA	7090
GAAACCACAG	CCCTGTGGTG	TGGGCCTCAC	ACCTTAGTGC	TCTGGTGGCC	TGATCTCCCA	GTGCCCTGCG	7160

FIGURE 9 (continued)

GGCAGCACAG	GATGTGGCTG	CTGGTGGAGG	TACCAACTGG	GCCCTGAACA	CAGGCCACAC	7230 ACCCCCATG
AGCCTGGGGA	CAGCATGAAA	AGTCTTATTT	GTTTCATGTG	ATATGATGTG	CCCTCACGAT	7300 TGCAGAGTGA
ACTCCACAAA	CTCTGAGGTA	ACTTGGGAAT	GTTCTTTTTT	TTTGAGACGG	AGTCTCACTC	7370 TGTCGCCCAG
GCTGGAGTGC	AGTGGCACAA	TCTTGGCTCA	CTGCAGCCTC	CACCTCCCAG	GTTCAAGTGA	7440 TTCTCCTGCC
TCAGCCCCC	AAATAGCTAG	GATTACAGGC	ACCGCCACCA	TGCCGGGCTA	ATTTTTTTGT	7510 ATTTTTAGTA
GAGATGGGGT	TTCACCATGT	TGGCCAGGCT	GGTTTGAAC	TCCTGACCTC	AAGTAATCCG	7580 CCCACCTCAG
CGTCCCAAAG	TGCTAGGATT	ACAGGCGTGA	GCCACCACTC	CCAACTGGGA	ATATTCTTGG	7650 GCACCGCACC
CATGGGAGCA	TGAAGGGTGG	ATGCAATGCA	ATCATAACAG	AGGCCCAAGG	TCAGCACTGG	7720 GGTGCTTGCC
TGTCATCCCA	GTGCTTTGGG	AGGCCGAGGT	GAGTGGATCG	TTAGAGCCCA	GGAGGTTGAG	7790 ACCAGCCTGG
GCAACATGGC	GAAACTCCCT	CTCTACAAAA	AGATACAAAA	ATTAGCCAGG	CAAGGTGGTG	7860 CACACCTGTA
GTCCAGCTA	CTCAGGAGAC	TGAGGTGGGA	GAATTGCCTG	AGCCTGGGGA	GGTCGAGGCT	7930 GCACTGACCT
GTGATCACAC	CACCACACTC	CAGCCTGGGT	GACAGTGAGA	CTCTGCCTCA	AAAAAACAAA	8000 AAATGAAAAA
ACCAGAGGCC	TCAGCCAATG	CCTGGGGGGC	TCAGAGTGCA	GCTGGCCCTT	CAGACGCTGA	8070 ACCAGTCATC
GGTAAAGGTT	TCCTCCAGGG	GCAGGAGGTG	TCCCAGTGGG	CAACAGTTCC	CCTCTGCCTA	8140 GCGTGATTCC
TGGGAAGGGA	CTCAGCTCAG	AGCCAACTCC	ACGTAGCTGG	AAATAAGGAC	CTCTGACCGA	8210 CTGGGGGTAG
GGTGGGTCT	GGGGTGGATC	CCTGCCCCAC	CCCCACAGCA	TCCCTACAGG	CATATCCTAC	8280 AGGCCTCGAA
GGTGCCTGGC	ACGTGGTGAG	AATGGTGCCA	GCGGCTGACC	CTGGCAGAGG	GCCAGGACTT	8350 GTCTCCAGCA
CCCATGTGCG	TGTTGCTTTA	TCCTTGCACT	GATCCACAGT	GGTAGCCACT	GATATTACCT	8420 TCATTTTACA
GATAGGGACA	CTGAAGTCCA	GAGAAGTTAA	GTAATGTGCC	TGAATTCACC	AATTAGCACG	8490 TGGCTGAGCT

FIGURE 9 (continued)

TATATTATTT	GTAGGGCTTC	AAAACACATG	GGAAATGGTT	TGTAAATCCA	AAATAATTCC	AAAATAAAGT	8560
TTATTAAAAC	TGAAAACAAT	ATGGCTTGGT	GTGGTGGCTC	ACACCTGTAA	TCCCAGCACT	TTGGGAGGCT	9960
NGAGGTGGGA	GTATTGCTTG	AGGCCAAGAG	TTCGAGACCA	GCCTGGGCAA	CATAGTGAGA	CCTTGCTCTT	10030
ACCAAAAACA	AAACAAAACA	AAAAACAAAG	CCAGGCATGT	GACGTGTGCC	TGTAGTTCCA	GCTACTTGGA	10100
GGCTGAGGCA	GGAGGATCAC	TTGAGGCCAG	GAGTTTGAGA	GACCCTGTCT	CTACAAAAAA	TTAAAATAAA	10170
AACAATAGTA	ACAGGCACTG	AGCCCTGGGC	CCTCCCCACT	GGCCTTTGCA	GTTTGCCTG	ATGCAGTACT	10240
					F A L	M Q Y	
CAAACTCCT	GAAGATCCAC	TTCACCTTCA	CCCAATTCCG	GACCAAGCCG	AGCCAGCAGA	GCCTGCTGGA	10310
S N L L	K I H	F T F T	Q F R	T S P	S Q Q S	L V D	
TCCCATCCTC	CAACTGAAAG	GCTTGACGTT	CACGSCCACG	GGCATCTGA	CAGTGCTGTA	AAGCAACCCC	10380
P I V	Q L K	G L T F	T A T	G I L T	V		

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